PCT/US03/08611

SEQUENCE LISTING

<110>	Transmolecular, GONDA, Matthew A GREENWOOD, John		40	508905
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4

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8

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Leu Tyr Gly Asp Ile Pro Pro Glu Leu Val Ala Lys Pro Leu Glu Asp 65 70 75 80

Leu Asp Pro Phe Tyr Lys Asp His Lys Thr Phe Met Val Leu Asn Lys 85 90 95

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12

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Asn Asp Ile Phe Arg Asn Leu Gln Lys Thr Val Ser Pro Lys Lys Gln 980 985 990

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Thr	Asp 1190		Asn	Met		Leu 1195		Phe	Thr	Glu	Val 1200		Asn	Arg
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Ser	Arg 1250		Lys	Asp	Glu	Gln 1255		Asp	Phe	Glu	Ala 1260		Leu	Tyr
	Tyr 1265		Tyr	Phe	Val	Val 1270	Phe	Ile	Ile	Phe	Gly 1275	Ser	Phe	Phe
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Ser	Met 1520		Cys	Leu	Phe	Gln 1525		Thr	Thr	Ser	Ala 1530		Trp	Asp
Thr 	Leu _ 1535	Leu 	Asn	Pro	Met 	Leu -1540	Glu	Ala	r̄ā=	_Glu	His_ 1545		_Asn	_Ser
Ser	Ser 1550		Asp	Ser	Сув	Gln 1555		Pro	Gln	Ile	Ala 1560		Val	Tyr

Phe	Val 1565	Ser	Tyr	Ile	Ile	Ile 1570	Ser	Phe	Leu	Ile	Val 1575	Val	Asn	Met
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Ser	Glu 1595	Asp	Pro	Leu	Gly	Glu 1600	Asp	Asp	Phe	Glu	Ile 1605	Phe	Tyr	Glu
Val	Trp 1610	Glu	Lys	Phe	Asp	Pro 1615	Glu	Ala	Ser	Gln	Phe 1620	IÌe	Gln	Tyr
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Lys	Leu 1700		Glu	Pro	Ile	Val 1705	Thr	Thr	Thr	Lys	Arg 1710	Lys	Glu	Glu
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678

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Ile Val Ile Gly Ile Ala Ile Val Ser Tyr Ile Pro Gly Ile Thr Ile

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	tca Ser													774
	gtg Val 250													822
	atc Ile													870
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	ttc Phe													1206
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18

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445

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			g gat ccc cga aag n Asp Pro Arg Ly: 980	
			c acc att gat ct r Thr Ile Asp Le 5	
			ccc aaa aag caa Pro Lys Lys Gln :	
			tgc ttt cca tgc Cys Phe Pro Cys	
agc gtg gac aag Ser Val Asp Lys	aga aag cct o Arg Lys Pro I 1035	ccc tgg gtc Pro Trp Val 1040	att tgg tgg aac Ile Trp Trp Asn	ctg 3165 Leu 1045
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					ata Ile 1215											3705
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									aag Lys 1535				agc Ser			4650
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Gln Pro Arg Pro Gln Leu Asp Leu Lys Ala Ser Arg Lys Leu Pro Lys 50 60

Leu Tyr Gly Asp Ile Pro Arg Glu Leu Ile Gly Lys Pro Leu Glu Asp 65 70 75 80

Leu Asp Pro Phe Tyr Arg Asn His Lys Thr Phe Met Val Leu Asn Arg 85 90 95

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Lys Ile Leu Ala Arg Gly Phe Ile Leu Asp Glu Phe Ser Phe Leu Arg 180 185 190

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Ser Tyr Ile Pro Gly Ile Thr Ile Lys Leu Pro Leu Arg Thr Phe 210 215 220

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Lys Arg Lys Ser Phe Phe Leu Arg Glu Ser Gly Lys Asp Gln Pro Pro Gly Ser Asp Ser Asp Glu Asp Cys Gln Lys Lys Pro Gln Leu Leu Glu Gln Thr Lys Arg Leu Ser Gln Asn Leu Ser Leu Asp His Phe Asp Glu His Gly Asp Pro Leu Gln Arg Gln Arg Ala Leu Ser Ala Val Ser Ile Leu Thr Ile Thr Met Lys Glu Gln Glu Lys Ser Gln Glu Pro Cys Leu Pro Cys Gly Glu Asn Leu Ala Ser Lys Tyr Leu Val Trp Asn Cys Cys Pro Gln Trp Leu Cys Val Lys Lys Val Leu Arg Thr Val Met Thr Asp Pro Phe Thr Glu Leu Ala Ile Thr Ile Cys Ile Ile Asn Thr Val Phe Leu Ala Met Glu His His Lys Met Glu Ala Ser Phe Glu Lys Met Leu Asn Ile Gly Asn Leu Val Phe Thr Ser Ile Phe Ile Ala Glu Met Cys Leu Lys Ile Ile Ala Leu Asp Pro Tyr His Tyr Phe Arg Arg Gly Trp Asn Ile Phe Asp Ser Ile Val Ala Leu Leu Ser Phe Ala Asp Val Met Asn Cys Val Leu Gln Lys Arg Ser Trp Pro Phe Leu Arg Ser Phe Arg Val Leu Arg Val Phe Lys Leu Ala Lys Ser Trp Pro Thr Leu-Asn _______67.5_. _ _ 68.0 Thr Leu Ile Lys Ile Ile Gly Asn Ser Val Gly Ala Leu Gly Ser Leu

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Asn	Pro	Thr	Gly 740	Pro	Thr	Val	Ser	Cys 745	Leu	Arg	His	Trp	His 750	Met	Gly
Asp	Phe	Trp 755	His	Ser	Phe	Leu	Val 760	Val.	Phe	Arg	Ile	Leu 765	Сув	Gly	Glu
Trp	Ile 770	Glu	Asn	Met	Trp	Glu 775	Сув	Met	Gln	Glu	Ala 780	Asn	Ala	Ser	Ser
Ser 785	Leu	Сув	Val	Ile	Val 790	Phe	Ile	Leu	Ile	Thr 795	Val	Ile	Gly	Lys	Leu 800
Val	Val	Leu	Asn	Leu 805	Phe	Ile	Ala	Leu	Leu 810	Leu	Asn	Ser	Phe	Ser 815	Asn
Glu	Glu	Arg	Asn 820	Gly	Asn	Leu	Glu	Gly 825	Glu	Ala	Arg	Lys	Thr 830	Lys	Val
Gln	Leu	Ala 835	Leu	Asp	Arg	Phe	Arg 840	Arg	Ala	Phe	Cys	Phe 845	Val	Arg	His
Thr	Leu 850	Glu	His	Phe	Cys	His 855	Lys	Trp	Cys	Arg	Lys 860	Gln	Asn	Leu	Pro
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Ile	Pro	Leu	Val	Met 885	Glu	Met	ГÀЗ	Arg	Gly 890		Glu	Thr	Gln	Glu 895	Glu
Leu	Gly	Ile	Leu 900	Thr	Ser	Val	Pro	Lys 905	Thr	Leu	Gly	Val	Arg 910	His	Asp
Trp	Thr	Trp 915			Pro		Ala 920	Glu 	Glu	Glu	Asp	Asp 925	Val	Glu	Ph∈

Ser Gly Glu Asp Asn Ala Gln Arg Ile Thr Gln Pro Glu Pro Glu Gln 930 935 940

'Gln Ala Tyr Glu Leu His Gln Glu Asn Lys Lys Pro Thr Ser Gln Arg

955 960 950 945

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Thr Ile Gln Asp Pro Arg Lys Lys Ser Asp Val Thr Ser Ile Leu Ser 980 985

Glu Cys Ser Thr Ile Asp Leu Gln Asp Gly Phe Gly Trp Leu Pro Glu 1000 995

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Gly Cys Cys Phe Pro Cys Cys Ser Val Asp Lys Arg Lys Pro Pro 1025 1030 1035

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Pro Lys Ile Gln Glu Leu Leu Asn Cys Thr Asp Ile Ile Phe Thr 1095 1085 1090

His Ile Phe Ile Leu Glu Met Val Leu Lys Trp Val Ala Phe Gly 1100 1105 1110

Phe Gly Lys Tyr Phe Thr Ser Ala Trp Cys Cys Leu Asp Phe Ile 1115 · 1120 1125

Ile Val Ile Val Ser Val Thr Thr Leu Ile Asn Leu Met Glu Leu 1130 1135 1140

Lys Ser Phe Arg Thr Leu Arg Ala Leu Arg Pro Leu Arg Ala Leu 1145 1150

_____ Ser Gln Phe Glu Gly Met Lys Val Val Val Asn Ala Leu Ile Gly 1165

Ala Ile Pro Ala Ile Leu Asn Val Leu Leu Val Cys Leu Ile Phe 1185 1180

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Phe	Gly 1205	Lys	Cys	Ile	Asn	Gly 1210	Thr	Asp	Ser	Val	Ile 1215	Asn	Tyr	Thr
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Ile	Asn 1235		Lys	Val	Asn	Phe 1240		Asn	Val	Gly	Asn 1245	Ala	Tyr	Leu
Ala	Leu 1250		Gln	Val	Ala	Thr 1255		ГХв	Gly	Trp	Met 1260	Asp	Ile	Ile
Tyr	Ala 1265		Val	Asp	Ser	Thr 1270		Lys	Glu	Gln	Gln 1275		Glu	Phe
Glu	Ser 1280	Asn	Ser	Leu	Gly	Tyr 1285	Ile	Tyr	Phe	Val	Val 1290	Phe	Ile	Ile
Phe	Gly 1295		Phe	Phe	Thr	Leu 1300	Asn	Leu	Phe	Ile	Gly 1305	Val	Ile	Ile
Asp	Asn 1310	Phe	Asn	Gln	Gln	Gln 1315	Lys	Lys	Leu	Gly	Gly 1320	Gln	Asp	Ile
Phe	Met 1325		Glu	Glu	Gln	Lys 1330		Tyr	Tyr	Asn	Ala 1335	Met	Lys	Lys
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Lys	Cys 1355		Gly	Leu	Val	Phe 1360	Aap	Ile	Val	Thr	Ser 1365	Gln	Ile	Phe
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Met	Ala 1385		Ser	-Tyr	-Asn	Gln 1390	Pro	ŗĀ	Ala	Met	Lys 1395	Ser	Ile	Leu
Asp	His 1400		Asn	Trp	Val	Phe 1405		Val	Ile	Phe	Thr 1410	Leu	Glu	Cys

Leu Ile Lys Ile Phe Ala Leu Arg Gln Tyr Tyr Phe Thr Asn Gly Trp Asn Leu Phe Asp Cys Val Val Leu Leu Ser Ile Val Ser Thr Met Ile Ser Thr Leu Glu Asn Gln Glu His Ile Pro Phe Pro Pro Thr Leu Phe Arg Ile Val Arg Leu Ala Arg Ile Gly Arg Ile Leu Arg Leu Val Arg Ala Ala Arg Gly Ile Arg Thr Leu Leu Phe 1475 1480 Ala Leu Met Met Ser Leu Pro Ser Leu Phe Asn Ile Gly Leu Leu 1495 1500 Leu Phe Leu Ile Met Phe Ile Tyr Ala Ile Leu Gly Met Asn Trp Phe Ser Lys Val Asn Pro Glu Ser Gly Ile Asp Asp Ile Phe Asn Phe Lys Thr Phe Ala Ser Ser Met Leu Cys Leu Phe Gln Ile Ser Thr Ser Ala Gly Trp Asp Ser Leu Leu Ser Pro Met Leu Arg Ser Lys Glu Ser Cys Asn Ser Ser Ser Glu Asn Cys His Leu Pro Gly Ile Ala Thr Ser Tyr Phe Val Ser Tyr Ile Ile Ile Ser Phe Leu 1580 1585 Ile Val Val Asn Met Tyr Ile Ala Val Ile Leu Glu Asn Phe Asn 1600 1605 Thr Ala Thr Glu Glu Ser Glu Asp Pro Leu Gly Glu Asp Asp Phe 1610 1620 Asp Ile Phe Tyr Glu Val Trp Glu Lys Phe Asp Pro Glu Ala Thr

Gln Phe Ile Lys Tyr Ser Ala Leu Ser Asp Phe Ala Asp Ala Leu 1645 Pro Glu Pro Leu Arg Val Ala Lys Pro Asn Lys Tyr Gln Phe Leu 1660 Val Met Asp Leu Pro Met Val Ser Glu Asp Arg Leu His Cys Met 1675 Asp Ile Leu Phe Ala Phe Thr Ala Arg Val Leu Gly Gly Ser Asp 1690 Gly Leu Asp Ser Met Lys Ala Met Met Glu Glu Lys Phe Met Glu 1700 1705 1710 Ala Asn Pro Leu Lys Lys Leu Tyr Glu Pro Ile Val Thr Thr Thr 1715 ' 1720 1725 Lys Arg Lys Glu Glu Glu Arg Gly Ala Ala Ile Ile Gln Lys Ala 1730 1735 1740 Phe Arg Lys Tyr Met Met Lys Val Thr Lys Gly Asp Gln Gly Asp 1745 1750 1755 Gln Asn Asp Leu Glu Asn Gly Pro His Ser Pro Leu Gln Thr Leu 1760 1765 1770 Cys Asn Gly Asp Leu Ser Ser Phe Gly Val Ala Lys Gly Lys Val 1775 1780 1785 His Cys Asp 1790 <210> 5 <211> 5822 <212> DNA <213> Mus musculus <220> <221> CDS <222> (19)..(5313) ₹220>

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							atc Ile 100					339
							aat Asn					387
							agc Ser					435
							aat Asn					483
							gtc Val					531
							aga Arg 180					579
							tgg Trp					627
-							ggt Gly					675
							aga Arg					723

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														atc Ile		819
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														tgc Cys		915
														tgg Trp		963
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Val 380	Val	Ile	Phe	Leu	Gly 385	Ser	Phe	Tyr	Leu	Leu 390	Asn	Leu	Thr	ctg Leu	Ala 395	1203
Val	Val	Thr	Met	Ala 400	Tyr	Glu	Glu	Gln	Asn 405	Arg	Asn	Val	Ala	gcc Ala 410	Glu	1251
Thr	Glu	Ala	Lys 415	Glu	Lys	Met	Phe	Gln 420	Glu	Ala	Gln	Gln	Leu 425	ttg Leu	Arg	1299
Glu	Glu	Lys 430	Glu	Āla	Leu	Val	Ala 435	Met	Gly	Ile	Āsp	Arg 440	Thr	tcc Ser	Leu	1347
Asn 	Ser .445	Leu 	Gln 	Ala	Ser	Ser 450	Phe	Ser	Pro	гÃã	Lys 455	Arg	_Ly.s.	.Phe.		1395
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cga	gcc	tca	gcg	tcc	gat	tca	gag	gac	gat	gcc	tct	aaa	aac	cca	caa	1491

Arg	Ala	Ser	Ala	Ser 480	Asp	Ser	Glu	Asp	Asp 485	Ala	Ser	Lys	Asn	Pro 490	Gln	
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			cac His													1587
			tta Leu													1635
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			cct Pro													1731
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gtt tcc ccc caa aag caa cca gat cga tgc ttt ccc aag ggc ctc Val Ser Pro Gln Lys Gln Pro Asp Arg Cys Phe Pro Lys Gly Lev 990 995 1000	c agt 3027 u Ser
tgt atc ttt cta tgt tgc aaa aca atc aaa aaa aag tcc ccc t Cys Ile Phe Leu Cys Cys Lys Thr Ile Lys Lys Lys Ser Pro 1 1005 1010 1015	tgg 3072 Trp
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Gln Pro Arg Pro Gln Leu Asp Leu Lys Ala Ser Arg Lys Leu Pro Lys 50 55 60

Leu Tyr Gly Asp Val Pro Pro Asp Leu Ile Ala Lys Pro Leu Glu Asp 65 70 75 80

Leu Asp Pro Phe Tyr Lys Asp His Lys Thr Phe Met Val Leu Asn Lys 85 90 95

Lys Arg Thr Ile Tyr Arg Phe Ser Ala Lys Arg Ala Leu Phe Ile Leu 100 105 110

Gly Pro Phe Asn Pro Ile Arg Ser Phe Met Ile Arg Ile Ser Val His 115 120 125

Ser Val Phe Ser Met Phe Ile Ile Cys Thr Val Ile Ile Asn Cys Met 130 135 140

Phe Met Ala Asn Asn Ser Ser Val Asp Ser Arg Pro Ser Ser Asn Ile 145 150 155 160

Pro Glu Tyr Val Phe Ile Gly Ile Tyr Val Leu Glu Ala Val Ile Lys 165 170 175

Ile Leu Ala Arg Gly Phe Ile Val Asp Glu Phe Ser Tyr Leu Arg Asp 180 185 190

Pro Trp Asn Trp Leu Asp Phe Ile Val Ile Gly Thr Ala Ile Ala Pro 195 200 205

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Val Ile Val Gly Ala Leu Leu Arg Ser Val Lys Lys Leu Val Asp Val

41

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Tyr Thr Asn Phe Asp Ser Phe Gly Trp Ser Phe Leu Ala Met Phe Arg 340 345 350

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Phe	Val 1565		Tyr	Ile	Ile	Ile 1570		Phe	Leu	Ile	Val 1575	Val	Asn	Met
Tyr	Ile 1580		Val	Ile	Leu	Glu 1585		Phe	Asn	Thr	Ala 1590		Glu	Glu
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 Arg Pro Phe Thr Ser Asp Ser Leu Ala Ala Ile Glu Lys Arg Ile Ala-----
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		gcg Ala														480
		ttc Phe														528
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		agc Ser														672
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		att Ile 275														864
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290 295 300

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Ile													ggc Gly			1152
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Tyr Leu Leu Asn Leu Thr Leu Ala Val Val Thr Met Ala Tyr Glu Glu 385 390 395 400

Gln Asn Arg Asn Val Ala Ala Glu Thr Glu Ala Lys Glu Lys Met Phe Gln Glu Ala Gln Gln Leu Leu Arg Glu Glu Lys Glu Ala Leu Val Ala Met Gly Ile Asp Arg Ser Ser Leu Asn Ser Leu Gln Ala Ser Ser Phe Ser Pro Lys Lys Arg Lys Phe Phe Gly Ser Lys Thr Arg Lys Ser Phe Phe Met Arg Gly Ser Lys Thr Ala Gln Ala Ser Ala Ser Asp Ser Glu Asp Asp Ala Ser Lys Asn Pro Gln Leu Leu Glu Gln Thr Lys Arg Leu Ser Gln Asn Leu Pro Val Asp Leu Phe Asp Glu His Val Asp Pro Leu His Arg Gln Arg Ala Leu Ser Ala Val Ser Ile Leu Thr Ile Thr Met Gln Glu Gln Glu Lys Phe Gln Glu Pro Cys Phe Pro Cys Gly Lys Asn Leu Ala Ser Lys Tyr Leu Val Trp Asp Cys Ser Pro Gln Trp Leu Cys Ile Lys Lys Val Leu Arg Thr Ile Met Thr Asp Pro Phe Thr Glu Leu Ala Ile Thr Ile Cys Ile Ile Ile Asn Thr Val Phe Leu Ala Val Glu His His Asn Met Asp Asp Asn Leu Lys Thr Ile Leu Lys Ile Gly Asn Trp Val Phe Thr Gly Ile Phe Ile Ala Glu Met Cys Leu Lys Ile Ile Ala Leu Asp Pro Tyr His Tyr Phe Arg His Gly Trp Asn Val Phe Asp

Ser Ile Val Ala Leu Leu Ser Leu Ala Asp Val Leu Tyr Asn Thr Leu Ser Asp Asn Asn Arg Ser Phe Leu Ala Ser Leu Arg Val Leu Arg Val 665 Phe Lys Leu Ala Lys Ser Trp Pro Thr Leu Asn Thr Leu Ile Lys Ile 675 680 Ile Gly His Ser Val Gly Ala Leu Gly Asn Leu Thr Val Val Leu Thr 690 695 Ile Val Val Phe Ile Phe Ser Val Val Gly Met Arg Leu Phe Gly Thr 705 710 Lys Phe Asn Lys Thr Ala Tyr Ala Thr Gln Glu Arg Pro Arg Arg Arg 725 Trp His Met Asp Asn Phe Tyr His Ser Phe Leu Val Val Phe Arg Ile 740 Leu Cys Gly Glu Trp Ile Glu Asn Met Trp Gly Cys Met Gln Asp Met 755 760 Asp Gly Ser Pro Leu Cys Ile Ile Val Phe Val Leu Ile Met Val Ile 770 775 Gly Lys Leu Val Val Leu Asn Leu Phe Ile Ala Leu Leu Leu Asn Ser 785 790 Phe Ser Asn Glu Glu Lys Asp Gly Ser Leu Glu Gly Glu Thr Arg Lys 805 815 Thr Lys Val Gln Leu Ala Leu Asp Arg Phe Arg Arg Ala Phe Ser Phe 820 825 830 Met Leu His Ala Leu Gln Ser Phe Cys Cys Lys Lys Cys Arg Arg Lys 835 840 Asn Ser Pro Lys Pro Lys Glu Thr Thr Glu Ser Phe Ala Gly Glu Asn _____850 ____ 850

Lys Asp Ser Ile Leu Pro Asp Ala Arg Pro Trp Lys Glu Tyr Asp Thr

865

Asp Met Ala Leu Tyr Thr Gly Gln Ala Gly Ala Pro Leu Ala Pro Leu 885 890 895

- Ala Glu Val Glu Asp Asp Val Glu Tyr Cys Gly Glu Gly Gly Ala Leu 900 905 910
- Pro Thr Ser Gln His Ser Ala Gly Val Gln Ala Gly Asp Leu Pro Pro 915 920 925
- Glu Thr Lys Gln Leu Thr Ser Pro Asp Asp Gln Gly Val Glu Met Glu 930 935 940
- Val Phe Ser Glu Glu Asp Leu His Leu Ser Ile Gln Ser Pro Arg Lys 945 950 955 960
- Lys Ser Asp Ala Val Ser Met Leu Ser Glu Cys Ser Thr Ile Asp Leu 965 970 975
- Asn Asp Ile Phe Arg Asn Leu Gln Lys Thr Val Ser Pro Lys Lys Gln 980 985 990
- Pro Asp Arg Cys Phe Pro Lys Gly Leu Ser Cys His Phe Leu Cys His 995 1000 1005
- Lys Thr Asp Lys Arg Lys Ser Pro Trp Val Leu Trp Trp Asn Ile 1010 1015 1020
- Arg Lys Thr Cys Tyr Gln Ile Val Lys His Ser Trp Phe Glu Ser 1025 1030 1035
- Phe Ile Ile Phe Val Ile Leu Leu Ser Ser Gly Ala Leu Ile Phe 1040 1045 1050
- Glu Asp Val Asn Leu Pro Ser Arg Pro Gln Val Glu Lys Leu Leu 1055 1060 1065
- Arg Cys Thr Asp Asn Ile Phe Thr Phe Ile Phe Leu Leu Glu Met 1070 1080
- Ala Trp Cys Trp Leu Asp Phe Leu Ile Val Val Ser Val Leu 1100 1105 1110
- Ser Leu Met Asn Leu Pro Ser Leu Lys Ser Phe Arg Thr Leu Arg

1115 1120 1125

Ala	Leu	Arg	Pro	Leu	Arg	Ala	Leu	Ser	Gln	Phe	Glu	Gly	Met	Lys
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- Val Val Val Tyr Ala Leu Ile Ser Ala Ile Pro Ala Ile Leu Asn 1145 1150 1155
- Val Leu Leu Val Cys Leu Ile Phe Trp Leu Val Phe Cys Ile Leu 1160 1165 1170
- Gly Val Asn Leu Phe Ser Gly Lys Phe Gly Arg Cys Ile Asn Gly 1175 1180 1185
- Thr Asp Ile Asn Met Tyr Leu Asp Phe Thr Glu Val Pro Asn Arg 1190 1195 1200
- Ser Gln Cys Asn Ile Ser Asn Tyr Ser Trp Lys Val Pro Gln Val 1205 1210 1215
- Asn Phe Asp Asn Val Gly Asn Ala Tyr Leu Ala Leu Leu Gln Val 1220 1235
- Ala Thr Tyr Lys Gly Trp Leu Glu Ile Met Asn Ala Ala Val Asp 1235 1240 1245
- Ser Arg Glu Lys Asp Glu Gln Pro Asp Phe Glu Ala Asn Leu Tyr 1250 1255 1260
- Ala Tyr Leu Tyr Phe Val Val Phe Ile Ile Phe Gly Ser Phe Phe 1265 1270 1275
- Thr Leu Asn Leu Phe Ile Gly Val Ile Ile Asp Asn Phe Asn Gln 1280 1285 1290
- Gln Gln Lys Lys Leu Gly Gly Gln Asp Ile Phe Met Thr Glu Glu 1295 1300 1305
- Gln Lys Lys Tyr Tyr Asn Ala Met Lys Lys Leu Gly Thr Lys Lys

 1310

 1315

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 - Pro Gln Lys Pro Ile Pro Arg Pro Leu Asn Lys Cys Gln Ala Phe 1325 1330 1335
 - Val Phe Asp Leu Val Thr Ser Gln Val Phe Asp Val Ile Ile Leu 1340 1345 1350

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Ala	Leu 1400	Arg	Gln	His	туг	Phe 1405	Thr	Asn	Gly	Trp	Asn 1410	Leu	Phe	Asp	
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Thr	Leu 1535	Leu	Asn	Pro	Met	Leu 1540	Glu	Ala	Lys	Glu	His 1545	Cys	Asn	Ser	
_ Ser.	Ser- 1550	Gln	-Asp	Ser-	Сув	Gln ⁻ 1555	-Gln	Pro	Gĺn	Īle	Ala 1560	Val	Val	Tyr	-
Phe	Val 1565	Ser	Tyr	Ile	Ile	Ile 1570	Ser	Phe	Leu	Ile	Val 1575	Val	Asn	Met	

Tyr Ile Ala Val Ile Leu Glu Asn Phe Asn Thr Ala Thr Glu Glu 1580 1585 1590 Ser Glu Asp Pro Leu Gly Glu Asp Asp Phe Glu Ile Phe Tyr Glu 1600 Val Trp Glu Lys Phe Asp Pro Glu Ala Ser Gln Phe Ile Gln Tyr 1610 1615 Ser Ala Leu Ser Asp Phe Ala Asp Ala Leu Pro Glu Pro Leu Arg 1625 1630 1635 Val Ala Lys Pro Asn Lys Phe Gln Phe Leu Val Met Asp Leu Pro 1645 1650 Met Val Met Gly Asp Arg Leu His Cys Met Asp Val Leu Phe Ala 1655 1660 1665 Phe Thr Thr Arg Val Leu Gly Asp Ser Ser Gly Leu Asp Thr Met 1670 1675 1680 Lys Thr Met Met Glu Glu Lys Phe Met Glu Ala Asn Pro Phe Lys 1685 1690 1695 Lys Leu Tyr Glu Pro Ile Val Thr Thr Lys Arg Lys Glu Glu 1700 1705 1710 Glu Gln Gly Ala Ala Val Ile Gln Arg Ala Tyr Arg Lys His Met 1715 1720 1725 Glu Lys Met Val Lys Leu Arg Leu Lys Asp Arg Ser Ser Ser Ser 1730 1740

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